

keywords: 932	Expectation: 3.0	UPE: 25.0	Search	Sort	miRNA Acc.
e.g. AT1G27360, miR156, transcription factor ...	Range: 0.0 - 3.0	Range: 0.0 - 25.0	by:	Expectation(E)	▼

List of Predicted miRNA/Target Pairs [#Session ID: 1332363080486463]

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miRNA Acc.	Target Acc.	Expectation (E)	Target Accessibility (UPE)	Alignment	Target Description	Inhibition	Multiplicity
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5461	Contig3553	2.0	15.553	miRNA 20 UCUUUUUUAUUAUGUGGAGA 1 ::: ::::: ::::: ::::: Target 16 AGAAAUAAAUAUACAUUUCU 35	Translation	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5461	LUSTC1NG- RP-155_B11_12MAR2007_093	3.0	8.486	miRNA 20 UCUUUUUUAUUAUGUGGAGA 1 ::: ::::: ::::: ::::: Target 27 AGAACUAAUGUAUCACCUUU 46	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5461	LUSFL1AD_Contig183	3.0	11.747	miRNA 20 UCUUUUUUAUUAUGUGGAGA 1 : :: ::::: ::::: Target 21 AUAAUUAGUAUACAUCUCU 40	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5461	LUSFL1AD_Contig157	3.0	12.747	miRNA 20 UCUUUUUUAUUAUGUGGAGA 1 : :: ::::: ::::: Target 114 AUAAUUAGUAUACAUCUCU 133	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5461	LUSGE1NG- RP-117_G04_09MAR2007_020	3.0	19.6	miRNA 20 UCUUUUUUAUUAUGUGGAGA 1 : ::::: ::::: Target 4 AAAAAUAAAUAUACAUUUCU 23	Translation	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig8500	2.0	6.989	miRNA 21 AAGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 729 UUUUUUUAAGAAAUAUAAUAA 749	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig5942	2.5	10.865	miRNA 20 AGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 51 UUUUUCAAGAAAAAAUAU 70	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig11328	2.5	18.478	miRNA 20 AGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 268 UCUUUGAGGAUAGAUAG 287	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig2943	3.0	12.051	miRNA 20 AGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 215 UCUUUUAAGUAACAAUAGA 234	Translation	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig7627	3.0	13.726	miRNA 20 AGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 465 UUCUCAAGUAUAGAUUA 484	Translation	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	LUSGC1NG- RP-140_D06_30JAN2007_042	3.0	15.888	miRNA 20 AGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 144 UCWUGAAAAGUAAUAGA 163	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	Contig13452	3.0	17.028	miRNA 21 AAGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 118 UGCUUUCAUCAAUAUAA 138	Cleavage	1	
scaffold1389.Lu0910.pre- miRNA:932.miRNA:5464	LUSTE1AD- RP-256_M20_9MAY2008_068	3.0	17.173	miRNA 21 AAGAAAGUUCUUUAUUAU 1 ::: ::::: ::::: Target 129 UGCUUUCAUCAAUAUAA 149	Cleavage	1	

scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig8500	2.0	6.814	miRNA Target	20 AAGAAAGUUCUUUAUUAU 1 ::: ::::::::::::::: 729 UUUUUUUAGAAAUAAUA 748	Cleavage 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig13467	3.0	10.437	miRNA Target	21 UAAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 25 AUUCUUUUAGAU AUGAUAUU 45	Translation 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig5942	3.0	10.721	miRNA Target	20 AAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 50 CUUUUUCAAGAAAAAUUA 69	Cleavage 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	LUSTE1AD-RP-256 M20 9MAY2008_068	3.0	16.754	miRNA Target	21 UAAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 128 GUGCUUUCAUCAAAUUAUA 148	Translation 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig13452	3.0	16.839	miRNA Target	21 UAAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 117 GUGCUUUCAUCAAAUUAUA 137	Translation 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig2784	3.0	17.838	miRNA Target	20 AAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 277 UUCUUUCAAGAACUGAUGUU 296	Cleavage 1
scaffold1389.Lu0910.pre-miRNA:932.miRNA:5465	Contig11328	3.0	18.263	miRNA Target	21 UAAGAAAGUUCUUUAUUAU 1 ::: ::::::: ::::::: 266 ACUCUUUUGAGGAUGAUUA 286	Cleavage 1